```
%% Specify Crystal and Specimen Symmetries
clear all
close all
CS = {'notIndexed',...
 crystalSymmetry('622', [3.2 3.2 5.2], 'X||a', 'Y||b*', 'Z||c', 'mineral', 'Mg', ✓
'color', 'light blue')};
% plotting conventions
setMTEXpref('xAxisDirection','east');
setMTEXpref('zAxisDirection','intoPlane');
%% Specify File Names
% which files to be imported
% fname = [pname '\0.08strain.ang'];0.08strain minimalCleanup.ang
fname = ['map20170109123448657.ang'];
%% Import the Data
% create an EBSD variable containing the data
ebsd = loadEBSD(fname,CS,'interface','ang',...
  'convertEuler2SpatialReferenceFrame');
ebsd(ebsd.prop.iq==0).phaseId=1;
ebsd=ebsd(ebsd.prop.ci>0.1);
oM = ipdfHSVOrientationMapping(ebsd('indexed'));
color = oM.orientation2color(ebsd('indexed').orientations);
seg angle = 15;
% minimum indexed points per grain between 5 and 10
min points = 5;
[grains,ebsd.grainId,ebsd.mis2mean] = calcGrains(ebsd('indexed'),...
    'angle', seg angle*degree, 'boundary', 'tight');
% remove small grains with less than min points indexed points
figure; plot (grains ('Mg'), grains ('Mg').meanOrientation)
grainsSelected = grains(grains.grainSize > 5);
%reindex ebsd
ebsd = ebsd(grainsSelected);
% figure;plot(grains('Mg'),grains('Mg').meanOrientation)
% and perform grain reconstruction with the reduces EBSD data set
[grains,ebsd.grainId,ebsd.mis2mean] = calcGrains(ebsd('indexed'),'angle',5*degree);
figure; plot (grains ('Mg'), grains ('Mg').meanOrientation)
%% Get pairs for twin relationships using mean orientation
```

```
tol=8*degree; %needs to be slight larger (like Rod noted) because dealing with\checkmark
meanOrientations
% We are going to use pairs to store all pairs information.
[counts,pairsTmp] = neighbors(grains);
pairs=struct;
pairs.type=zeros(length(pairs),1,'int8');
pairs.combine=zeros(length(pairs),1,'logical');
pairs.pairs=pairsTmp
pairs.grainId1=pairsTmp(:,1);
pairs.grainId2=pairsTmp(:,2);
pairs.len=length(pairsTmp);
%Specify the twin definitions
tt1 = orientation('axis', Miller(1, -2, 1, 0, ebsd{2}.CS, 'uvw'),...
  'angle',86.3471*degree,ebsd{2}.CS,ebsd{2}.CS); %Extension twin <10-11> largest amount
tt2 = orientation('axis', Miller(1, 0, -1, 0, ebsd{2}.CS, 'uvw'), ...
  'angle',34.7*degree,ebsd{2}.CS,ebsd{2}.CS); %Extension twin <1126> Quite a few
ct1 = orientation('axis', Miller(1, 0, -1, 0, ebsd{2}.CS, 'uvw'), ...
  'angle',64.3*degree,ebsd{2}.CS,ebsd{2}.CS); %Main contraction twin
%check if pairs contain twins
for i=1:pairs.len
    mori=inv(grains(pairs.grainId1(i)).meanOrientation)*grains(pairs.grainId2(i)). ✓
meanOrientation; %Compute misorientation between all pairs
    [pairs.combine(i),pairs.type(i)] = TestTwinRelationship(ebsd,mori,tol,tt1,tt2,ct1);
end
pairs.grainId1 TC=pairs.grainId1(pairs.combine)
pairs.grainId2 TC=pairs.grainId2(pairs.combine)
pairs.type TC=pairs.type(pairs.combine)
pairs.pairs TC=pairs.pairs(pairs.combine,1:2)
grains.prop.boundaryType=pairs.type
% grains.prop.type(pairs(combine,1:2))
% figure;plot(grains(pairs(combine,1:2)))
\$\$ Compute schmid factor for the various twin combinations (Complete) Moving to in grainm{arksigma}
calculations since I care about pairs and mapping is a pain.
CRSS TT1=111;
CRSS TT2=113;
CRSS CT1=195;
sigma = tensor([0 0 0; 0 0 0; 0 0 -1], 'name', 'stress') %Sign of loading does more than <math>\checkmark
just invert twin/parent flag
subgrains1 = grains(pairs.grainId1); %could just pass pairs
subgrains2 = grains(pairs.grainId2);
subgrains1.prop.type=pairs.type;
subgrains2.prop.type=pairs.type;
[sFRelative12,sF1,sF2,SFActiveVar] = GetSchmidRelative(subgrains1,subgrains2,tt1,tt2,ct1, 🗸
CRSS TT1, CRSS TT2, CRSS CT1, sigma);
```

```
%sF is stored for pairs where 1 and 2 are for each. Right now not computing
%none twin sF
pairs.sF1=sF1';
pairs.sF2=sF2';
pairs.sFRelative12=sFRelative12';
pairs.sFActiveVar=SFActiveVar;
pairs.sF1 TC=pairs.sF1(pairs.combine);
pairs.sF2 TC=pairs.sF2(pairs.combine);
grains.prop.sF1=pairs.sF1;
grains.prop.sF2=pairs.sF2;
grains.prop.sFRelative12=pairs.sFRelative12;
grains.prop.sFActiveVar=pairs.sFActiveVar;
%plot schmid relationships. relationships can plot over each other because not grouped
%yet.
% plotType=2;
% figure
% plot(subgrains1(subgrains1.prop.type==plotType),pairs.sF1 TC(pairs.type TC==plotType))
% plot(subgrains2(subgrains1.prop.type==plotType),pairs.sF2 TC(pairs.type TC==plotType))
% hold off
%% Construct intergranular graph
s=pairs.pairs(:,1)';
t=pairs.pairs(:,2)';
G=graph(s,t);
%Comments: Once we have all the pertinent data in the graph we no longer
%have to worry about indexing and mapping between grains and merged grains
%etc... This is the primary reason that I chose the graph method!
% Add node labels and other grain level information
G.Nodes.Name=cellstr(int2str(grains.id))
G.Nodes.Id=str2num(cell2mat(G.Nodes.Name));
G.Nodes.Area=grains.area;
G. Nodes. Perimeter=grains.perimeter;
G.Nodes.AspectRatio=grains.aspectRatio;
G.Nodes.Paris=grains.calcParis;
G.Nodes.centroids=grains.centroid;
G.Nodes.meanOrientation=grains.meanOrientation
G.Nodes.Properties.UserData.mineral=grains.mineral; %For single phase material
% Compute grain boundary for each grain fragment (i.e. node)
G.Nodes.Gb=cell(length(grains),1);
for i=1:length(grains)
    G.Nodes.Gb{i}=grains(i).boundary;
end
% Add intergranular information
G.Edges.pairs=pairs.pairs;
G.Edges.type=pairs.type'; %Twin relation type
```

```
G.Edges.SF=[pairs.sF1,pairs.sF2]; %Schmid factors for relation type
G.Edges.SFRelative12=pairs.sFRelative12;
%Want to filter out SFRelative values that come from schmid factors with same sign?
%use the voting scheme
% G.Edges.SFRelative12(sum(sign(G.Edges.SF(:,1:2)),2)==2)=0
G.Edges.SFActiveVar=pairs.sFActiveVar';
% G.Edges.disorientation=
% Make a list of boundaries shared between two grains connected by an edge
G.Edges.Gb=cell(pairs.len,1);
G.Edges.ebsdId=cell(pairs.len,1);
grain1=grains(pairs.grainId1);
grain2=grains(pairs.grainId2);
for i=1:length(grain1)
    id1=grain1(i).boundary.ebsdId;
    id2=grain2(i).boundary.ebsdId;
    [boundaryEbsdId, loc] = intersect (id1, id2, 'rows');
    G.Edges.Gb{i}=grain1.boundary(loc);
    G.Edges.ebsdId{i}=boundaryEbsdId;
end
%Overlayer graph on grains
figure;
h=plot(grains, grains.meanOrientation, 'Micronbar', 'off')
set(gca, 'Units', 'normalized');
savefig('grains.fig');
hold on
p=plot(G,'XData',G.Nodes.centroids(:,1),'YData',G.Nodes.centroids(:,2))
hold off
p.Marker='s';p.NodeColor='k';p.MarkerSize=3;p.EdgeColor='k';
%% Remove nodes that have grains that are too small
% G=rmnode(G,grains(grains.grainSize < 5).id)</pre>
%% Reduce intergranular graph to grain clusters
%(i.e. delete edges between neighboring grain clusters)
%Remove edges that don't satisfy the twin relationship
G.Edges.Removed=G.Edges.type==0;
G clust=rmedge(G,G.Edges.pairs(G.Edges.Removed,1),...
    G.Edges.pairs(G.Edges.Removed,2))
%Using the gplot gui editor to delete unwanted connections
% A=adjacency(G clust);
% xy=G clust.Nodes.centroids;
% graph gui(A,xy,grains)
% Overlayer graph on grains
figure;
```

```
plot(grains, grains.meanOrientation)
hold on
p=plot(G clust,'XData',G clust.Nodes.centroids(:,1),'YData',G clust.Nodes.centroids(:,2))
p.Marker='s';p.NodeColor='k';p.MarkerSize=3;p.EdgeColor='k';
labeledge(p,1:length(G clust.Edges.pairs),1:length(G clust.Edges.pairs))
%% Remove edges from list generated visually (improve on this later
edgeList2Remove=[]; % number of with respect to the graph above
G clustClean=rmedge(G clust,edgeList2Remove)
figure;
plot(grains, grains.meanOrientation)
p=plot(G clustClean, 'XData', G clustClean.Nodes.centroids(:,1), 'YData', G clustClean.Nodes. ✓
centroids(:,2))
hold off
p.Marker='s';p.NodeColor='k';p.MarkerSize=3;p.EdgeColor='k';
labeledge(p,1:length(G clustClean.Edges.pairs),1:length(G clustClean.Edges.pairs))
%% Identify Families in grain clusters
%Remove nodes that aren't connected by edges, alternatively use condition
%on minimum occurance from conncomp
Grains2Keep=unique(G clustClean.Edges.pairs)
G clustClean=rmnode(G clustClean,G clustClean.Nodes.Id(~ismember(G clustClean.Nodes.Id, ✓
Grains2Keep)));
G clustClean.Nodes.Group = conncomp(G clustClean)'; %Get the nodes that are connected by \checkmark
edges
%Asign each pair to a cluster of grains
G clustClean.Edges.Group=zeros(length(G clustClean.Edges.pairs),1);
for i=1:length(G clustClean.Edges.pairs)
      G clustClean.Edges.Group(i)=G clustClean.Nodes.Group(...
          find(G_clustClean.Edges.pairs(i,1)==G_clustClean.Nodes.Id));
end
%loop over grain clusters to determine parent, grain groups
G clustClean.Nodes.FamilyID=zeros(length(G clustClean.Nodes.Id),1);
for i=1:max(G clustClean.Edges.Group)
    egroupId= find((i==G clustClean.Edges.Group)==true); %convert logical arrays to 🗸
indices
    ngroupId= find((i==G clustClean.Nodes.Group)==true);
    ori=G clustClean.Nodes.meanOrientation(ngroupId);
    G clustClean.Nodes.FamilyID(ngroupId) = GetFamily(ori)
end
%Determine what family each pair relates
G clustClean.Edges.FamilyID=zeros(length(G clustClean.Edges.pairs),2);
```

```
for i=1:max(G clustClean.Edges.Group)
    egroupId= find((i==G clustClean.Edges.Group)==true); %convert logical arrays to ✓
indices
   ngroupId= find((i==G clustClean.Nodes.Group)==true);
    nId=G clustClean.Nodes.Id(ngroupId);
    fId=G clustClean.Nodes.FamilyID(ngroupId);
    for j=1:length(egroupId)
        G clustClean.Edges.FamilyID(egroupId(j),1)=unique(fId(G clustClean.Edges.pairs ✓
(egroupId(j),1) == nId));
        G clustClean.Edges.FamilyID(egroupId(j),2)=unique(fId(G clustClean.Edges.pairs ✓
(egroupId(j),2) == nId));
    end
end
plot(grains(G clustClean.Nodes.Id), G clustClean.Nodes.FamilyID, 'Micronbar', 'off')
p=plot(G clustClean, 'XData', G clustClean.Nodes.centroids(:,1), 'YData', G clustClean.Nodes. ✓
centroids(:,2))
hold off
p.Marker='s';p.NodeColor='k';p.MarkerSize=3;p.EdgeColor='k';
labeledge(p,1:length(G clustClean.Edges.pairs),1:length(G clustClean.Edges.pairs))
%% Implement voting scheme for the respective families
% For each family compute the vote between connected families
%Edges currently connect clustered grains. We can compute the vote for each
%edge and then average for the family. This ensures that only neighbors are
%ever voted on. So that votes are really between familys average values of
%schmid are computed and stored for each fragment in a family. The area and
%relative boundary are taken in the addative sense rather than average.
%Edges.Gb contains boundaries between pairs
%Nodes.Gb contains boundaries for each fragment
%To compute the relative boundary between families we need to sum boundary
% of a particular type and since each pair is its own type this should be
%relatively simple.
w = [1, 1, 1]
G clustClean.Nodes.Properties.UserData.Mineral=grains.mineral;
G Complete = FamilyVotes(G clustClean, w);
G_Complete.Nodes.isTwin=zeros(length(G_Complete.Nodes.Id),1);
G Complete.Nodes.isAParent=zeros(length(G Complete.Nodes.Id),1,'logical');
for i=1:max(G Complete.Edges.Group)
    egroupId= find((i==G Complete.Edges.Group)==true); %converts logical arrays to ✓
indices
    ngroupId= find((i==G Complete.Nodes.Group)==true);
    nFamily = G Complete.Nodes.FamilyID(ngroupId)
    nId = G Complete.Nodes.Id(ngroupId)
    eType = G Complete. Edges. type (egroupId)
```

```
eVote = G Complete.Edges.Vote(egroupId,:)
    ePairs = G Complete.Edges.pairs(egroupId,:)
    eFamily = G Complete.Edges.FamilyID(egroupId,:)
    %Initialize parent
    Parent = zeros(size(ePairs,1),2,'logical');
    %Make list of each family relation
    FamilyRelationList=cell(max(nFamily),1);
    for j=1:max(nFamily)
        FamilyInPair=ismember(eFamily(:,:),j)
        FamilyRelationList{j}=FamilyInPair
        FamilyRelationList{j}
        eFamily
    end
    %The number of relation types for each family
    %Here we also assign the parent for the case of two tensile variants
    %having the same family.
    numTypeFamily=zeros (max (nFamily), 1);
    for j=1:max(nFamily)
        id=logical(sum(ismember(eFamily(:,:),j),2))
        uniqueTypes=unique(eType(id))
        numTypeFamily(j)=length(uniqueTypes)
        if numTypeFamily(j)>1
            if ismember([1,2],uniqueTypes) %tensile twins can't make tensile twins
                Parent(FamilyRelationList{j})=true;
            end
        end
    end
    %Assign the rest of the parents
    for j=1:size(Parent,1)
        if sum(Parent(j,:))==0
            [\sim, loc] = max(eVote(j,:));
            Parent(j,loc) = true;
        end
    end
    %Finally assign twin label to nodes (need to differentiate generation)
    G Complete.Nodes.isTwin(ngroupId(ismember(nId,ePairs(~Parent))))=5; eType
    G Complete.Nodes.isAParent(ngroupId(ismember(nId,ePairs(Parent))))=true;
end
plot(grains(G Complete.Nodes.Id),G Complete.Nodes.isTwin,'Micronbar','off')
figure;
plot(grains(G Complete.Nodes.Id), grains(G Complete.Nodes.Id). ✓
meanOrientation,'Micronbar','off')
```

```
ori1=G_clustClean.Nodes.meanOrientation(pairId(j,1));
        ori2=G clustClean.Nodes.meanOrientation(pairId(j,1));
figure;
plot(grains(333),grains(333).meanOrientation,'Micronbar','off')
p=plot(G clustClean,'XData',G clustClean.Nodes.centroids(:,1),'YData',G clustClean.Nodes. 🗸
centroids(:,2))
hold off
p.Marker='s';p.NodeColor='k';p.MarkerSize=3;p.EdgeColor='k';
labeledge(p,1:length(G clustClean.Edges.pairs),1:length(G clustClean.Edges.pairs))
%% Now we want to merge the grains together into sets
%Build a list of boundaries to merge
gBList=G clustClean.Edges.Gb{1}('Mg','Mg');
for i=2:length(G clustClean.Edges.pairs)
    gBList=[BoundaryList;G clustClean.Edges.Gb{i}('Mg','Mg')];
end
[mergedGrains, parentId] = merge(grains, gBList('Mg', 'Mg'));
% copy ebsd data into a new variable to not change the old data
ebsd merged = ebsd;
% update the grainIds to the parentIds
ebsd merged('indexed').grainId = parentId(ebsd('indexed').grainId)
figure
plot(grains, grains.meanOrientation)
hold on
plot(mergedGrains.boundary,'k','linewidth',4)
plot(grains.boundary, 'linecolor', 'r', 'linewidth', 1)
text(grains,int2str(grains.id))
hold off
[MergedGrains, ParentId] = merge (grains, BoundaryList);
% copy ebsd data into a new variable to not change the old data
ebsd merged = ebsd;
% update the grainIds to the parentIds
ebsd merged('indexed').grainId = parentId(ebsd('indexed').grainId)
plot(grains, grains.meanOrientation)
hold on
plot(mergedGrains.boundary,'k','linewidth',4)
plot(grains.boundary, 'linecolor', 'r', 'linewidth', 1)
text(grains, int2str(grains.id))
hold off
%% Get boundaries between EBSD points
[twinBoundary_tt1,twinBoundary_tt2,twinBoundary_ct1,twinBoundary_sec1, \( \n' \)
twinBoundary ttwint] = GetTwinBoundaries(ebsd,grains.boundary('Mg','Mg'),5*degree);
combinedTwinTot=[twinBoundary tt1;twinBoundary tt2;twinBoundary ct1] %twinBoundary tt1; ✓
```

```
twinBoundary tt2; twinBoundary ct1;
%% Let's see how we have done
figure
plot(ebsd, ebsd.orientations)
hold on
% plot(gB('Mg','Mg'),'linecolor','k','linewidth',6)
% plot(combinedTwinTot,'linecolor','r','linewidth',2)
plot(twinBoundary tt1, 'linecolor', 'r', 'linewidth', 2)
plot(twinBoundary tt2, 'linecolor', 'b', 'linewidth', 2)
plot(twinBoundary ct1, 'linecolor', 'g', 'linewidth', 2)
% plot(twinBoundary sec1, 'linecolor', 'b', 'linewidth',2)
text(grains,int2str(grains.id))
% scatter(boundaryList(:,1),boundaryList(:,2))
hold off
%% Add and remove grain pairs that are not likely to be twins using previous graph
pairs.toRemove=[103,241;299,349;398,440;511,440;441,511;606,505;...
    470, 429; 290, 225; 312, 328; 682, 683; 658, 641; 410, 366; 636, 592; 649, 600; ...
    646,593;168,203;270,211;306,360;287,360;275,360;306,328;360,293;211,207;...
    304,306;308,3061;
pairs.toAdd=[];%168,203 Need to search for the pair in pairs.pairs. It won't have a ∠
schmid factor so set type and schmid factor to 0;
pairs.combineClean=pairs.combine;
pairs.typeClean=pairs.type;
%make all pair combinations and make sure they can't/should be combined
pairs.toRemove=[pairs.toRemove;fliplr(pairs.toRemove)];
[boundaryLoc,loc]=intersect(pairs.pairs.toRemove,'rows');
pairs.combineClean(loc)=false;
pairs.typeClean(loc)=0;
pairs.pairsClean=[pairs.pairs];%;pairs.toAdd];
pairs.combineClean=[pairs.combineClean];%;ones(size(pairs.toAdd,1),1,'logical')];
%Recalculate pairs for the modified data
pairs.pairs TCClean=pairs.pairsClean(pairs.combineClean,1:2);
pairs.grainId1 TCClean=pairs.pairsClean(pairs.combineClean,1);
pairs.grainId2 TCClean=pairs.pairsClean(pairs.combineClean,2);
pairs.type TCClean=pairs.typeClean(pairs.combineClean);
pairs.sF1 TC=pairs.sF1(pairs.combineClean);
pairs.sF2 TC=pairs.sF2(pairs.combineClean);
% Redo calculation from new pairs list
boundaryList=[];
grain1=grains(pairs.pairsClean(pairs.combineClean,1));
grain2=grains(pairs.pairsClean(pairs.combineClean, 2));
for i=1:length(grain1)
    id1=grain1(i).boundary.ebsdId;
    id2=grain2(i).boundary.ebsdId;
    [boundaryLoc, loc] = intersect(id1, id2, 'rows');
```

```
gBtmp=grain1.boundary(loc);
   boundaryList=[boundaryList;boundaryLoc];
end
id=grains.boundary.ebsdId;
[boundaryLoc,loc]=intersect(id,boundaryList,'rows');
gB_clean=grains.boundary(loc)
figure
plot(grains, grains.meanOrientation)
plot(gB clean('Mg','Mg'),'linecolor','k','linewidth',6)
% plot(combinedTwinTot,'linecolor','r','linewidth',2)
plot(twinBoundary tt1,'linecolor','r','linewidth',2)
plot(twinBoundary tt2, 'linecolor', 'b', 'linewidth', 2)
plot(twinBoundary ct1, 'linecolor', 'g', 'linewidth', 2)
% plot(twinBoundary sec1, 'linecolor', 'b', 'linewidth',2)
text(grains,int2str(grains.id))
% scatter(boundaryList(:,1),boundaryList(:,2))
hold off
%% Merge grains into clusters for twin/parent analysis
[mergedGrains, parentId] = merge(grains, gB clean('Mg', 'Mg'));
% copy ebsd data into a new variable to not change the old data
ebsd merged = ebsd;
% update the grainIds to the parentIds
ebsd merged('indexed').grainId = parentId(ebsd('indexed').grainId)
figure
plot(grains, grains.meanOrientation)
hold on
plot(mergedGrains.boundary,'k','linewidth',4)
plot(grains.boundary, 'linecolor', 'r', 'linewidth', 1)
text(grains,int2str(grains.id))
hold off
%Good to here. Now we need to segment the combined grains.
% From here on we should be dealing with the merged grain set?
%% Create Families
%plot the mergedGrains IDs
figure
plot(grains, grains.meanOrientation)
hold on
plot(mergedGrains.boundary, 'k', 'linewidth', 4)
plot(grains.boundary,'linecolor','r','linewidth',1)
text (mergedGrains, int2str (mergedGrains.id))
hold off
grains.prop.familyID=zeros(length(grains),1,'int8');
%For schmid calculations
```

```
CRSS TT1=111; %from cpfe
CRSS TT2=113;
CRSS CT1=195;
sigma = tensor([0 0 0; 0 0 0; 0 0 -1], 'name', 'stress') %Sign of loading does more than ✓
just invert twin/parent flag
grains.prop.familyID=zeros(length(grains),1,'int8');
grains.prop.sF1=zeros(length(grains),1,'int8');
grains.prop.sF2=zeros(length(grains),1,'int8');
grains.prop.grainId1=zeros(length(grains),1,'int8');
grains.prop.grainId2=zeros(length(grains),1,'int8');
% grains.prop.grainId1=grains(ismember(grains.id,pairs.grainId2)
groups=struct;
groups.len=length(mergedGrains);
groups.family=cell(groups.len,1)
groups.grains={}
% groups.combine=zeros(length(groups),1,'logical');
% groups.groups=groupsTmp
% groups.grainId1=groupsTmp(:,1);
% groups.grainId2=groupsTmp(:,2);
% groups.len=length(groupsTmp);
for i=184%1:length(mergedGrains)
    groups.grains{i}=grains(parentId==mergedGrains(i).id);
    [counts2,pairs2]=neighbors(groups.grains{i})
    figure;plot(groups.grains{i},groups.grains{i}.meanOrientation)
    %group similar orientations
    len=length(grainCluster);
    oriSame=zeros(len,len,'logical');
    FamiliesComplete=zeros(len,1,'logical');
    cnt=0;
    for j=1:len
        for k=1:len
            oriSame(j,k)=angle(grainCluster(j).meanOrientation,grainCluster(k). ✓
meanOrientation) /degree <10;</pre>
        end
        if FamiliesComplete(oriSame(j,:))~=true
            cnt=cnt+1;
            FamiliesComplete(oriSame(j,:))=true;
            grains.prop.familyID(grainCluster.id(oriSame(j,:)))=cnt;
        end
        if sum(FamiliesComplete) ==len
        elseif sum(FamiliesComplete) ==len-1
            cnt=cnt+1;
```

```
grains.prop.familyID(grainCluster.id(~FamiliesComplete))=cnt;
            break;
        end
    end
    %Compute grain family properties
    for j=1:cnt
        family=grains.prop.familyID(grainCluster.id) == j;
        grains.prop.familyGbLen(grainCluster.id(family))=grainCluster(family). <
boundarySize; %boundary lengths of families
        grains.prop.familySize(grainCluster.id(family))=grainCluster(family).area;
        %Add schmid here?
응
          [countsFamilies, pairsFamilies] = neighbors (grains (grainCluster.id));
          m(j)=mean(ebsd.orientations(ismember(ebsd.grainId,grains.id(grainCluster.id

✓
9
(family)))))
   end
        %get pairs inside merged grain
        id=(ismember(pairsClean(combineClean,1),grainCluster.id)+...
            ismember(pairsClean(combineClean,2),qrainCluster.id)); %PairsClean consists ✓
of grain ids
        pairID=find(id==2) %Should be the indices of pairs that are in our combined grain
        %Determine family pair and direction for each schmid pair
        pairFamily=cell(cnt,1);
        for j=1:cnt
            family=grains.prop.familyID(grainCluster.id)==j;
            pairFamily{j}=[ismember(pairsClean(combineClean(pairID),1),grainCluster.id

✓
(family)),...
                ismember(pairsClean(combineClean(pairID),2),grainCluster.id(family))]; % ✓
this indexes pairsClean
        end
        familyFs=grains.prop.sF1(ismember(grains.id,pairsClean(pairFamily{1},1)))
        grains.prop.sF2(ismember(grains.id,pairsClean(pairFamily{1},1)))
        grains.prop.sF2(ismember(grains.id,pairsClean(pairFamily{3},1)))
        grains.prop.sF2(pairID)
        intersect(pairFamily{1},pairFamily{2})
        subgrains1 = grains(pairs(combine,1));
        subgrains2 = grains(pairs(combine,2));
        subgrains1.prop.type=grains.prop.type(combine);
        subgrains2.prop.type=grains.prop.type(combine);
        [sFl,sF2] = GetSchmidRelative(subgrains1, subgrains2, tt1, tt2, ct1, CRSS TT1, ✓
CRSS TT2, CRSS CT1, sigma);
        grains.prop.sF1=zeros(length(grains),1,'int8');
        grains.prop.sF2=zeros(length(grains),1,'int8');
        grains.prop.sF1(combine) = sFl';
        grains.prop.sF2(combine) = sF2';
```

%plot schmid relationships. relationships can overright because not grouped

```
%yet.
        plotType=subgrains1.prop.type==1;
        figure
        plot(subgrains1(plotType), sFl(plotType))
        plot (subgrains2 (plotType), sF2 (plotType))
        hold off
          grains.prop.familySFtt1(grainCluster.id(family))=grains.prop.sF1(grainCluster. ✓
id(family))
          grains.prop.familySFtt2
양
          grains.prop.familySFct1
end
% end
% grains.prop.familyID(grainCluster.id)
plot(grains, grains.prop.familyID)
hold on
plot(mergedGrains.boundary, 'k', 'linewidth', 4)
hold off
%% Compute boundary lengths of families
%% Now we need to make a list of boundaries shared between the grains
boundaryList=[];
grain1=grains(pairs(combine,1));
grain2=grains(pairs(combine,2));
for i=1%:length(grain1)
   x1=grain1(i).boundary.x;
    y1=grain1(i).boundary.y;
    x2=grain2(i).boundary.x;
    y2=grain2(i).boundary.y;
    [boundaryLoc,loc]=intersect([x1,y1],[x2,y2],'rows');
    gB=grain1.boundary(loc);
    if i==1
          boundaryList=gB('Mg','Mg');
9
        boundaryList=boundaryLoc;
    else
응
          boundaryList=[boundaryList;gB('Mg','Mg')];
        boundaryList=[boundaryList;boundaryLoc];
    end
    % scatter(boundarLoc(:,1),boundarLoc(:,2))
     plot(grain1.boundary(loc),'k','linewidth',3)
     hold off
응
```

```
응
      [mergedGrains,parentId] = merge(grains,gB('Mg','Mg'));
end
%Since this clearly works
x=grains.boundary.x; y=grains.boundary.y;
[boundaryLoc, loc]=intersect([x,y], boundaryList, 'rows');
xy=[grains.boundary.x(loc),grains.boundary.y(loc)];
%works for xy but xy location index ~=boundary index WHY!!!!
%segmentIds are meaningless..
figure
plot(ebsd('Mg'),ebsd('Mg').orientations)
hold on
plot(gB,'k','linewidth',3)
% scatter(boundaryList(:,1),boundaryList(:,2))
scatter(xy(:,1),xy(:,2),'k')
hold off
응응
plot(grains)
hold on
plot(grains(pairs(163,1:2)), 'yellow')
hold off
[mergedGrains, parentId] = merge(grains, combinedTwin);
% copy ebsd data into a new variable to not change the old data
ebsd merged = ebsd;
% update the grainIds to the parentIds
ebsd merged('indexed').grainId = parentId(ebsd('indexed').grainId)
plot(ebsd merged('Mg'), ebsd('Mg').orientations)
plot(mergedGrains.boundary,'k','linewidth',3)
hold off
% hold on
% plot(grains2merge('Mg','Mg'),'linewidth',3)
% hold off
% for i=1:numGrains
     if ~ismember(i,processNeighbors)
응
양
          cnt=cnt+1;
          processNeighbors(cnt)=i;
응
      end
```

```
양
응
      while length(processNeighbors) <= cnt</pre>
          [counts,pairs] = neighbors(grains(grainId))
응
응
          group=zeros(counts,1,'logical');
양
          for j=1:counts
              mis=angle(grains(i).meanOrientation, grains(pairs(j,2)).meanOrientation) / 🗸
응
degree
양
              twinDected = TestTwinRelationship(ebsd, mis, tol)
양
              if twinDetected
              end
          end
% end
gB = grains.boundary
gB MgMg = gB('Mg','Mg')
% figure
plot(grains('Mg'), grains('Mg').meanOrientation)
% plot(ebsd('Mg'),ebsd('Mg').orientations)
hold on
% plot the boundary of all grains
plot(grains.boundary,'linewidth',4)
hold off
%% Get twin boundaries
[twinBoundary tt1, twinBoundary tt2, twinBoundary ct1, twinBoundary sec1, \( \mu \)
twinBoundary ttwint] = GetTwinBoundaries(ebsd,gB MgMg,5*degree);
% plot the twinning boundaries
% figure
plot(grains('Mg'),grains('Mg').id(randperm(length(grains('Mg')))))
% plot(ebsd,ebsd.orientations)
hold on
%plot(gB MgMg,angle(gB MgMg.misorientation,twinning),'linewidth',4)
plot(twinBoundary_tt1, 'linecolor', 'w', 'linewidth', 2, 'displayName', 'Tensile Twin 1 🗸
Boundary')
plot(twinBoundary tt2,'linecolor','w','linewidth',2,'displayName','Tensile Twin 2 ✓
Boundary')
plot(twinBoundary ct1,'linecolor','w','linewidth',2,'displayName','Compression Twin 1┟
Boundary')
plot(twinBoundary sec1, 'linecolor', 'w', 'linewidth', 2, 'displayName', 'Secondary Twin 1 ✓
Boundary')
plot(twinBoundary ttwint, 'linecolor', 'w', 'linewidth', 2, 'displayName', 'Tensile-Tensile ✓
Twin Boundary')
text(grains,int2str(grains.id))
hold off
legend off
```

```
%% Merge grains and label the parent grains
combinedTwinTot=[twinBoundary_tt1;twinBoundary_tt2;twinBoundary_sec1;twinBoundary_ttwint] 🗸
%; twinBoundary ct1 excluded because easy to identify
combinedTwin=combinedTwinTot;
%% Compute the misorientation between mean orientations to segment the microstructure
%% Need to remove twin boundary for merging that lies on parent grain boundaries.
numGrains=grains.length
ratio gBs=zeros(numGrains,1)
for i=1:numGrains
    %Check if is x,y against all boundaries
    %If a twin has a twin boundary relationship with
    grainIdGlobal=or(gB MgMg.grainId(:,1)==i,gB MgMg.grainId(:,2)==i);
    grainIdTwin=or(combinedTwin.grainId(:,1)==i,combinedTwin.grainId(:,2)==i);
    tmp gB=gB MgMg(grainIdGlobal);
    tmp gB isTwinning=combinedTwin(grainIdTwin);
    ratio gBs(i) = length(tmp gB isTwinning) / length(tmp gB);
    if ratio gBs(i) < 0.0 \&\& ratio <math>gBs(i) > 0
        combinedTwin(grainIdTwin) = [];
    end
end
% figure
% plot(grains)
% hold on
% % plot the boundary of all grains
% plot(,'linewidth',4)
% % plot(combinedTwin(grainIdTwin), 'linewidth', 4, 'linecolor', 'w')
% hold off
[mergedGrains, parentId] = merge(grains, combinedTwin);
% copy ebsd data into a new variable to not change the old data
ebsd merged = ebsd;
% update the grainIds to the parentIds
ebsd merged('indexed').grainId = parentId(ebsd('indexed').grainId)
figure
plot(ebsd merged('Mg'),ebsd('Mg').orientations)
plot(mergedGrains.boundary,'k','linewidth',3)
[twinBoundary_tt1,twinBoundary tt2,twinBoundary ct1,twinBoundary sec1, ✓
```

```
twinBoundary ttwint] = GetTwinBoundaries(ebsd, gB MgMg, 5*degree);
[mergedGrains,parentId] = merge(mergedGrains,combinedTwinTot);
% plot the merged grains
%plot(ebsd,ebsd.orientations)
figure
plot(mergedGrains.boundary, 'linecolor', 'k', 'linewidth', 2.5, 'linestyle', '-', ...
  'displayName', 'merged grains')
hold on
text (mergedGrains, int2str (mergedGrains.id))
plot(twinBoundary tt1,'linecolor','r','linewidth',2,'displayName','Tensile Twin 1')
plot(twinBoundary tt2, 'linecolor', 'g', 'linewidth', 2, 'displayName', 'Tensile Twin 2')
plot(twinBoundary ct1,'linecolor','b','linewidth',2,'displayName','Compression Twin')
plot(twinBoundary sec1, 'linecolor', 'y', 'linewidth', 2, 'displayName', 'Secondary Twin')
plot(twinBoundary_ttwint, 'linecolor', 'c', 'linewidth', 2, 'displayName', 'Tensile-Tensile ✓
Twin Boundary')
hold off
legend off
%% Loop over grains and pull twins out by their size in pixels
% gBInner=mergedGrains.innerBoundary;
% gBPar
% isTwinning = angle(gBInner.misorientation,tt1) < 7.5*degree;</pre>
% twinBoundaryInner=gBInner(isTwinning)
%Note: the parent ID maps each grain in original scan to a merged grain.
%We want to to extract all of the grains within a merged grain. Finally we
%Want to reform the parent grain and visualize the result to make sure the
%algorithm is doing what we want.
% Make grain id map and compare to IPF map
% figure
% plot(grains, grains.id(randperm(length(grains))), 'micronbar', 'off', 'figSize', 'large')
% plot(grains.boundary)
% mtexColorMap hsv
% hold off
% color = oM.orientation2color(ebsd.orientations);
% plot(ebsd, color, 'micronbar', 'off', 'figSize', 'large')
%from here we need to compute:
%1) should have check for
%1) twin boundary length for all boundary types: parent, neighbor, twin-twin
%2) group merged grains by misorientation
%3) grain area
%4) schmid factor at parent/twin boundary for twin type
%with this data in arrays we can:
%1) apply voting scheme to segment structure and assign twin types
%2) compute twin thickness
```

```
%3) varient type
%reconstruction strategy is
isTwin=zeros(length(grains),1);
nmerged=length (mergedGrains)
for i=1:nmerged
    subgrains = grains(parentId == mergedGrains(i).id);
    maxArea=max(subgrains.grainSize);
    % need to group similar same orientations into single parent grain so
    % we compute the misorientation between mean components
    flag=false; cnt=1;
    while cnt <=length(subgrains) && flag==false</pre>
        mis=angle(subgrains(cnt).meanOrientation, subgrains.meanOrientation)./degree;
        area=sum(subgrains.grainSize(mis<5));</pre>
        if area >=maxArea
            isTwin(subgrains(mis>5).id)=1;
            flag=true;
        end
        cnt=cnt+1;
    end
end
% figure
plot(grains, isTwin, 'micronbar', 'off', 'figSize', 'large')
hold on
% plot(grains.boundary)
plot(twinBoundary tt1, 'linecolor', 'w', 'linewidth', 2, 'displayName', 'Tensile Twin 1 ✓
Boundary')
hold off
```